

FIG. 1A

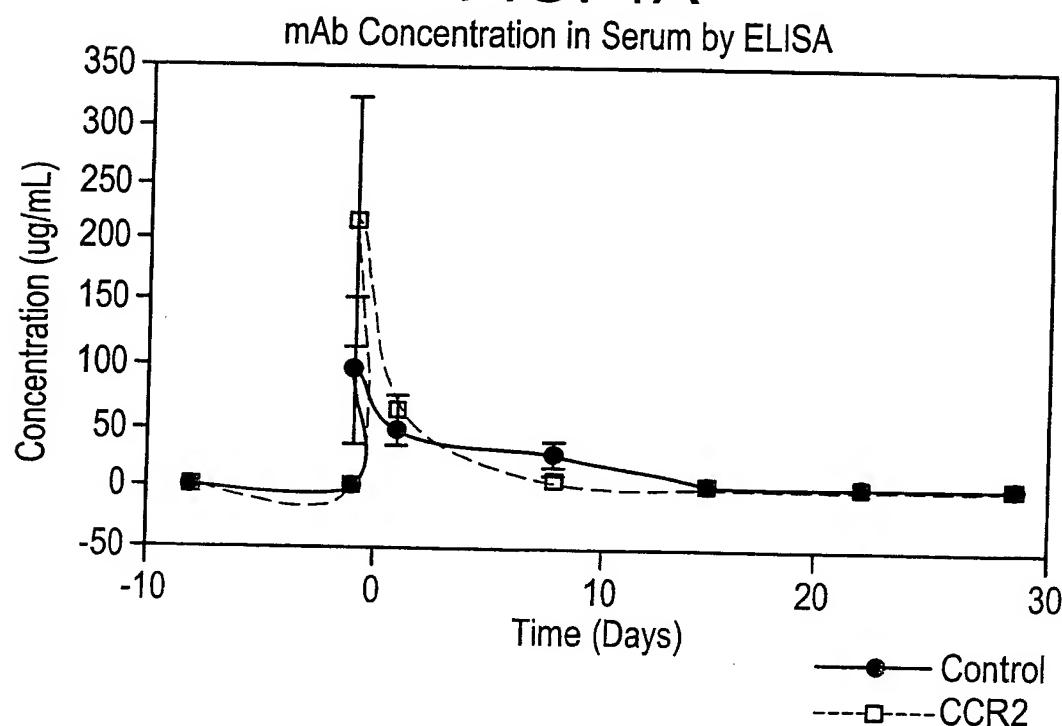


FIG. 1B

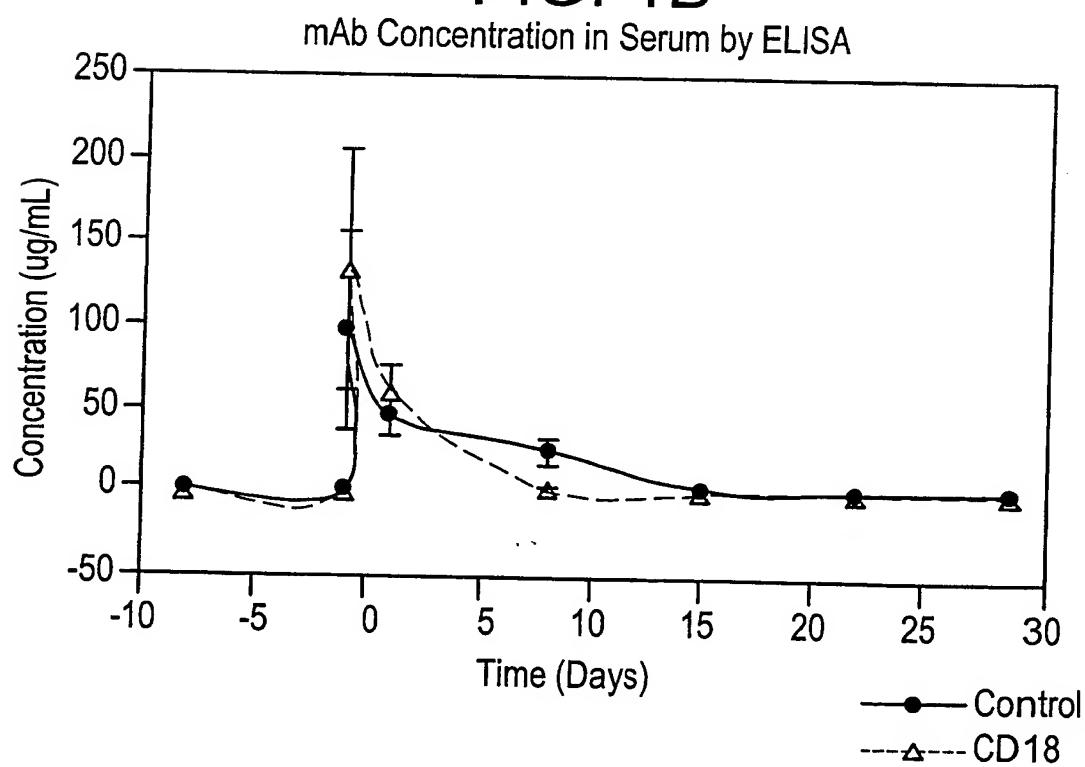


FIG. 2A

Monocyte Free Target Sites

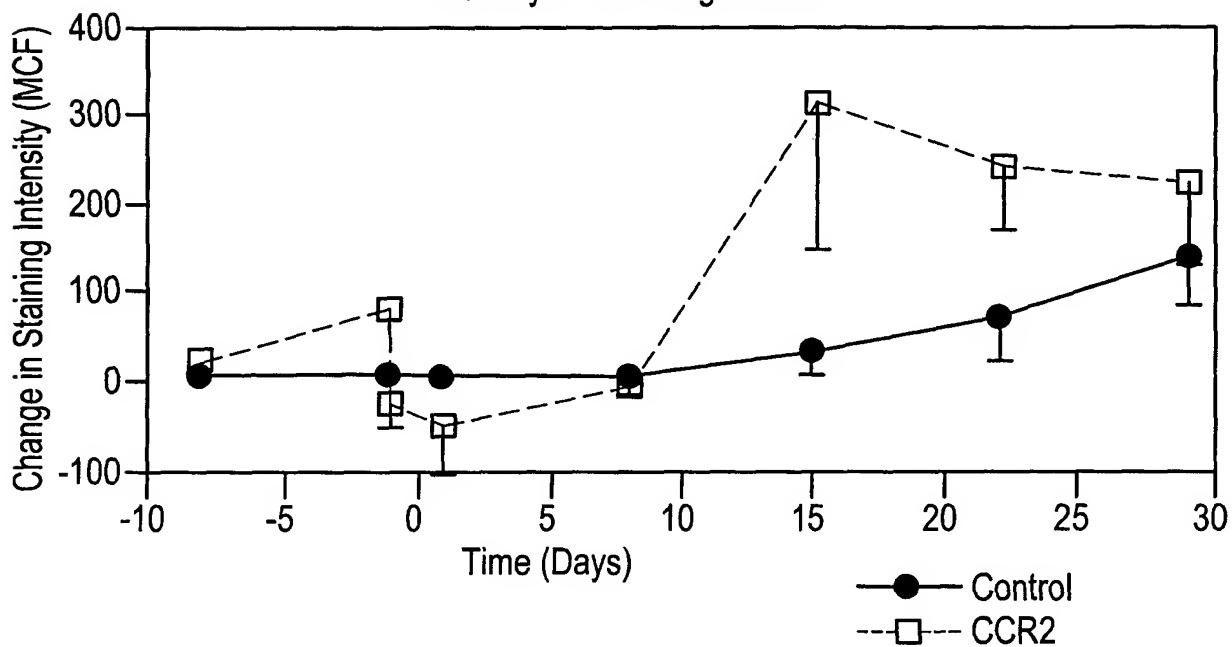


FIG. 2B

Neutrophil Free Target Sites

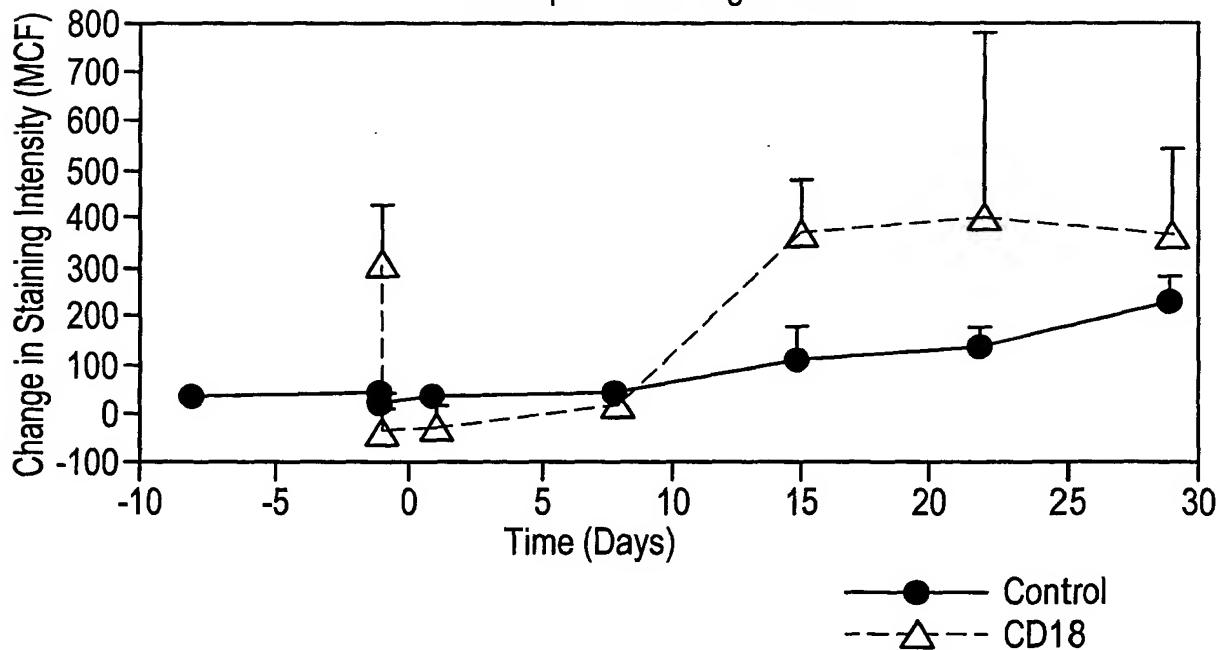


FIG. 2C

Monocyte Free Target Sites

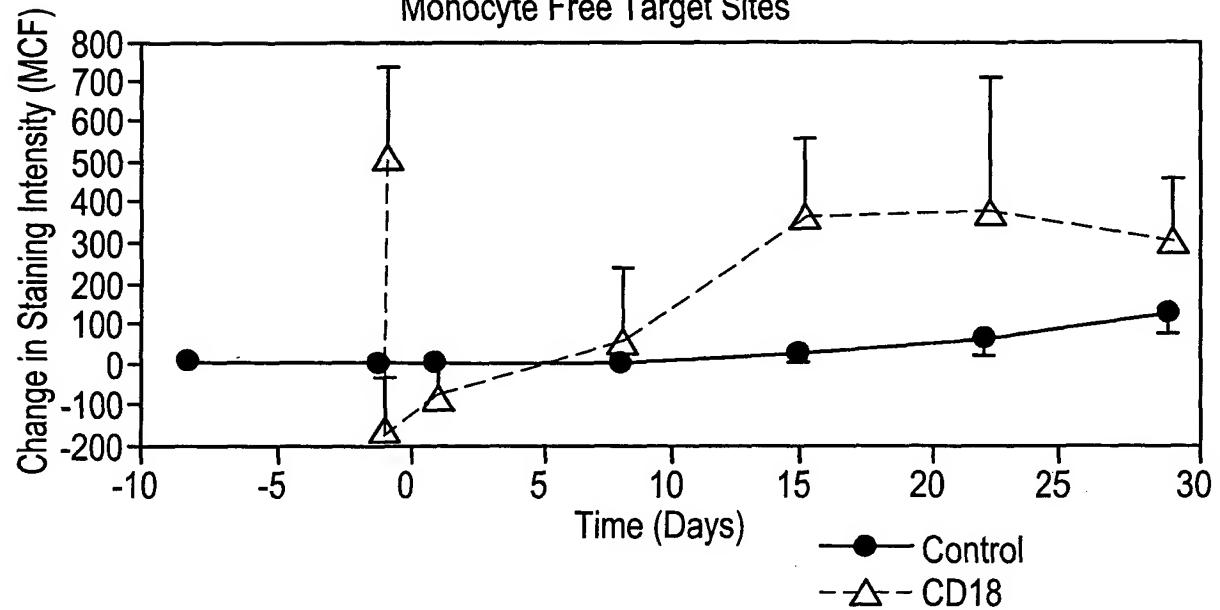


FIG. 3A

Total WBC Count

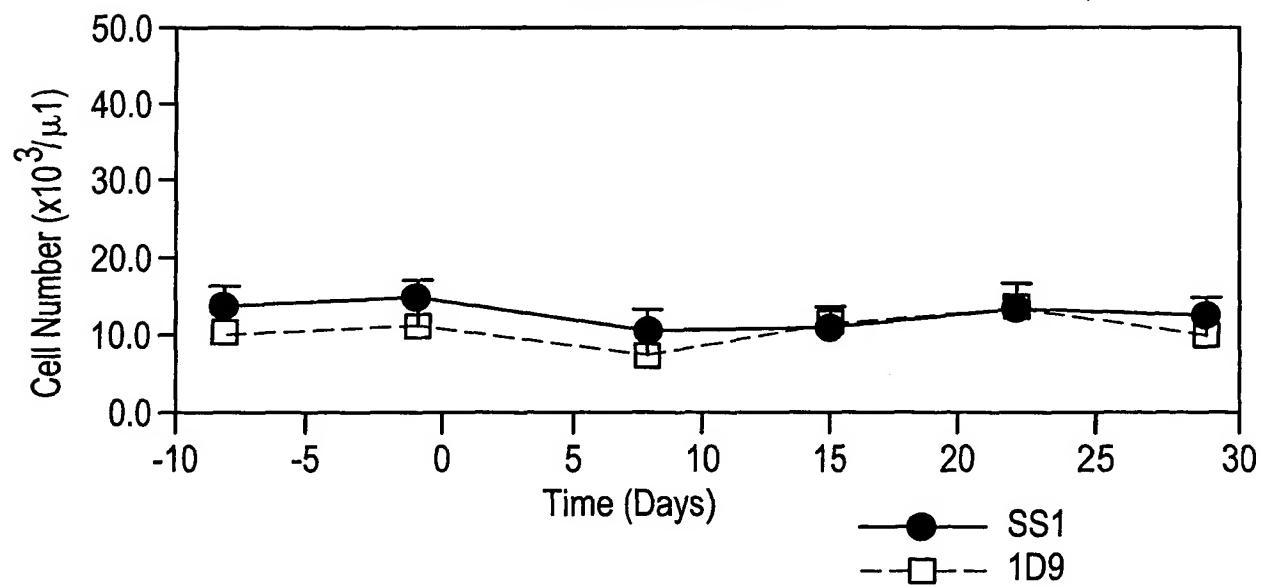


FIG. 3B

Total Neutrophil Count

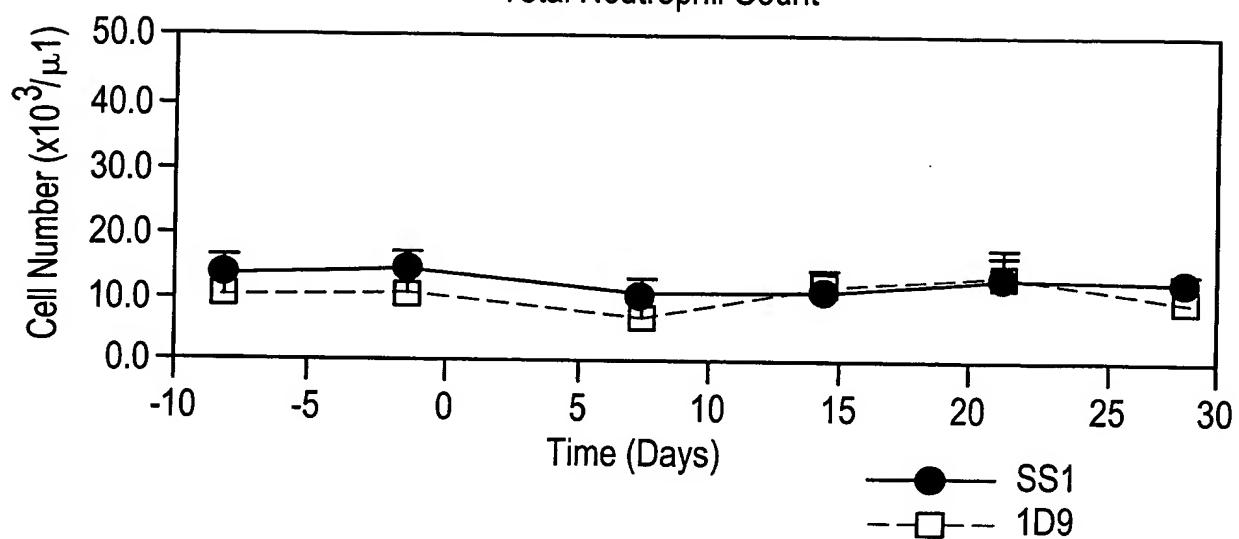


FIG. 3C

Total Lymphocyte Count

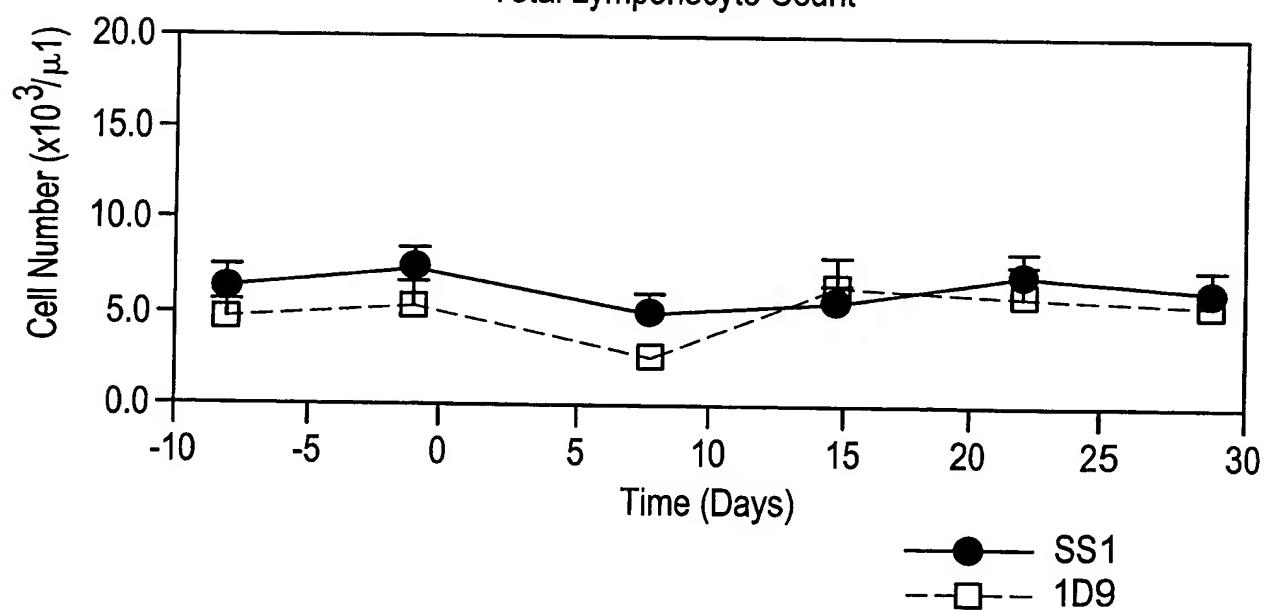


FIG. 3D
Total Monocyte Count

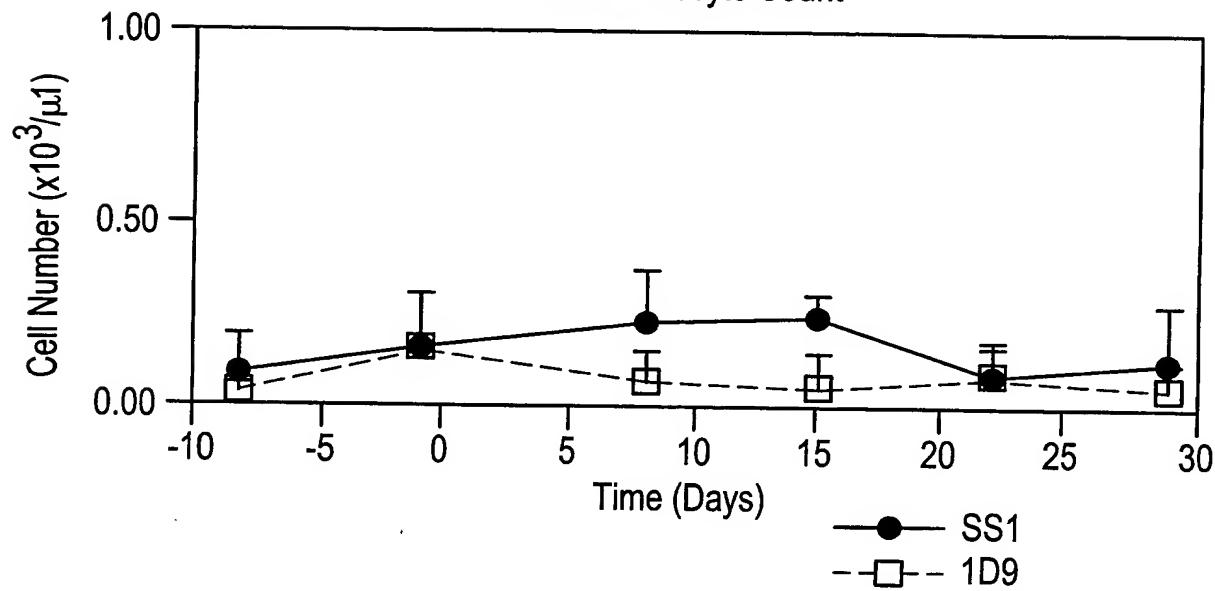


FIG. 3E

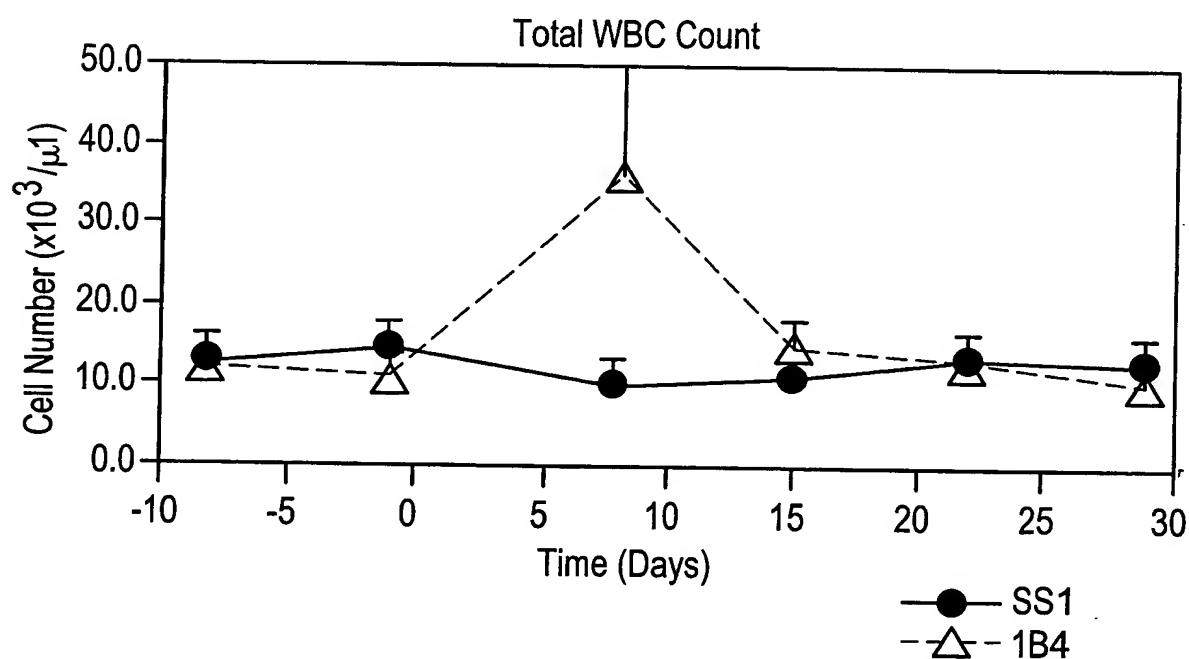


FIG. 3F

Total Neutrophil Count

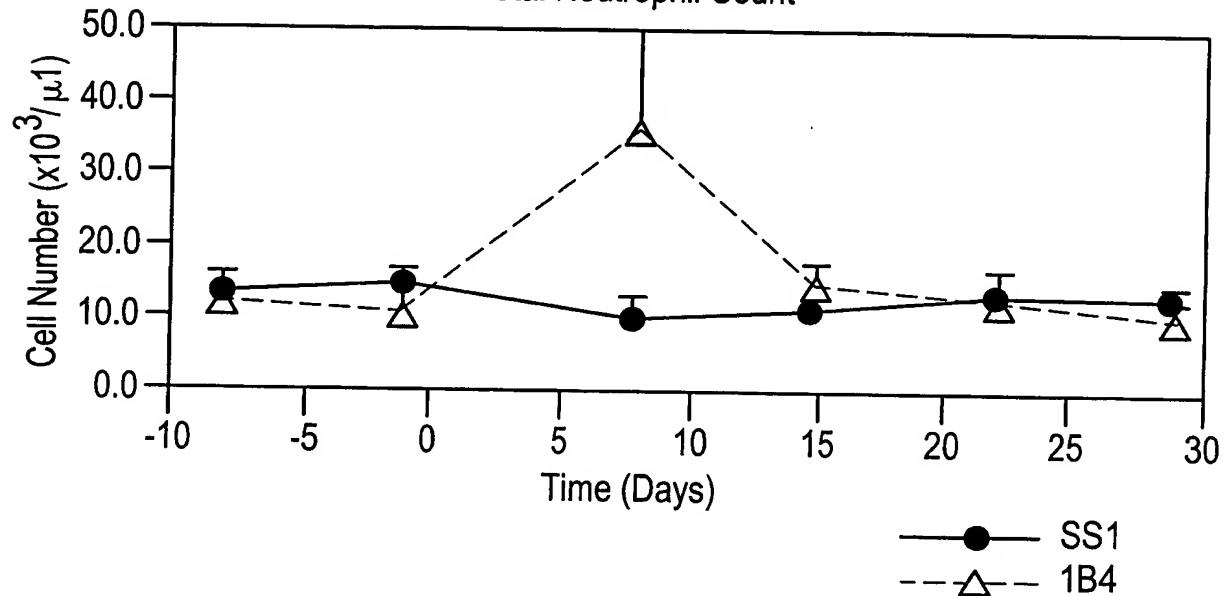


FIG. 3G

Total Lymphocyte Count

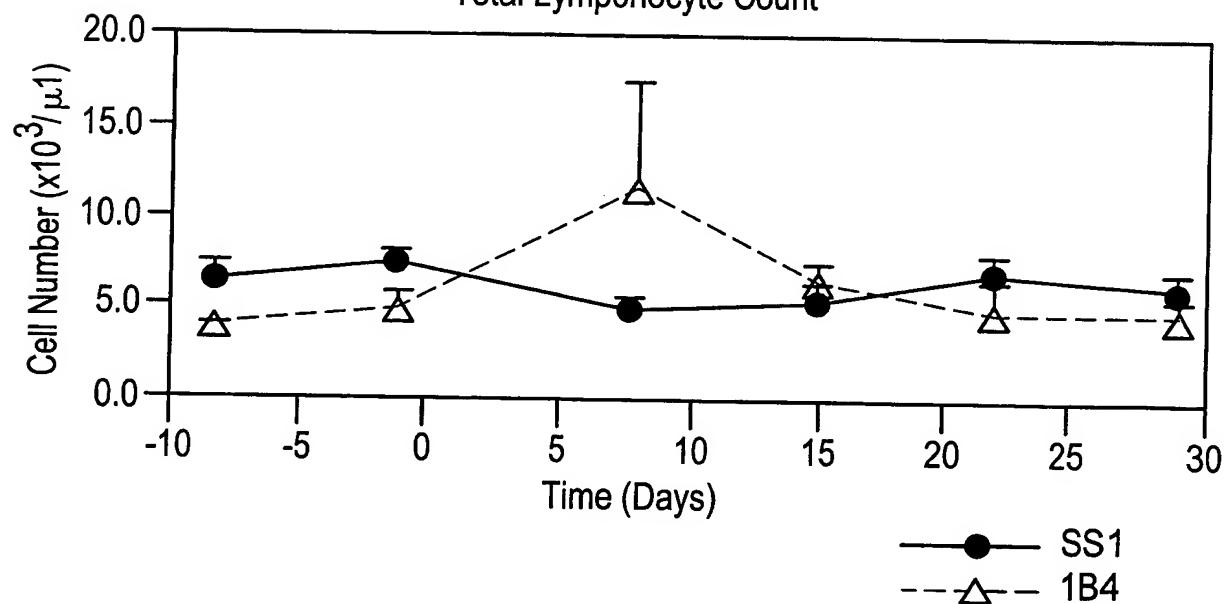


FIG. 3H
Total Monocyte Count

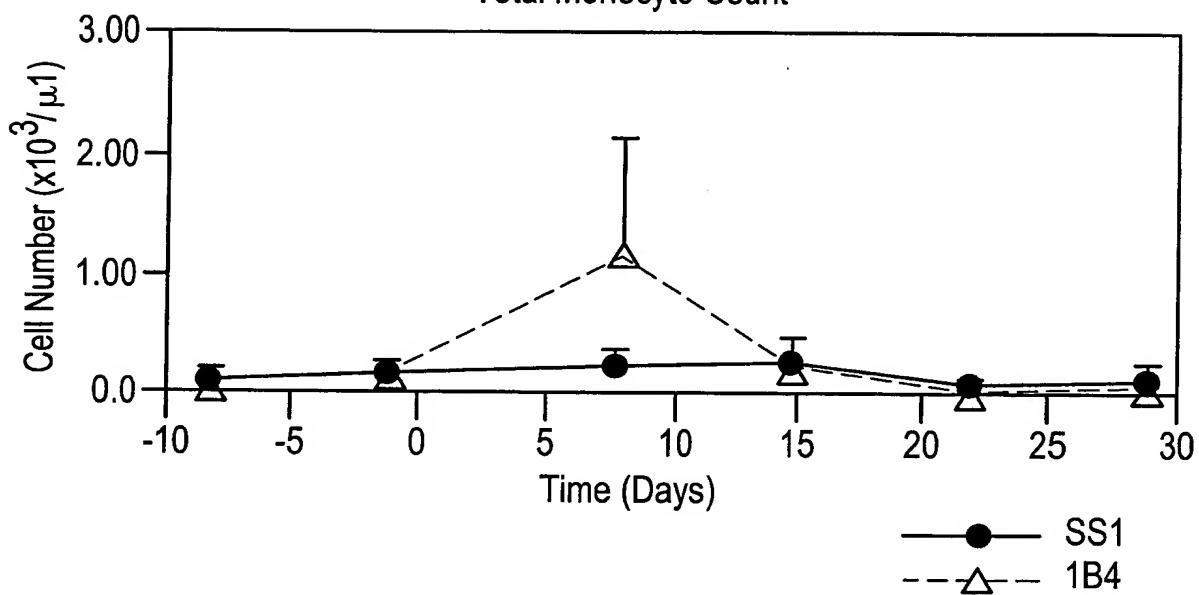


FIG. 4A

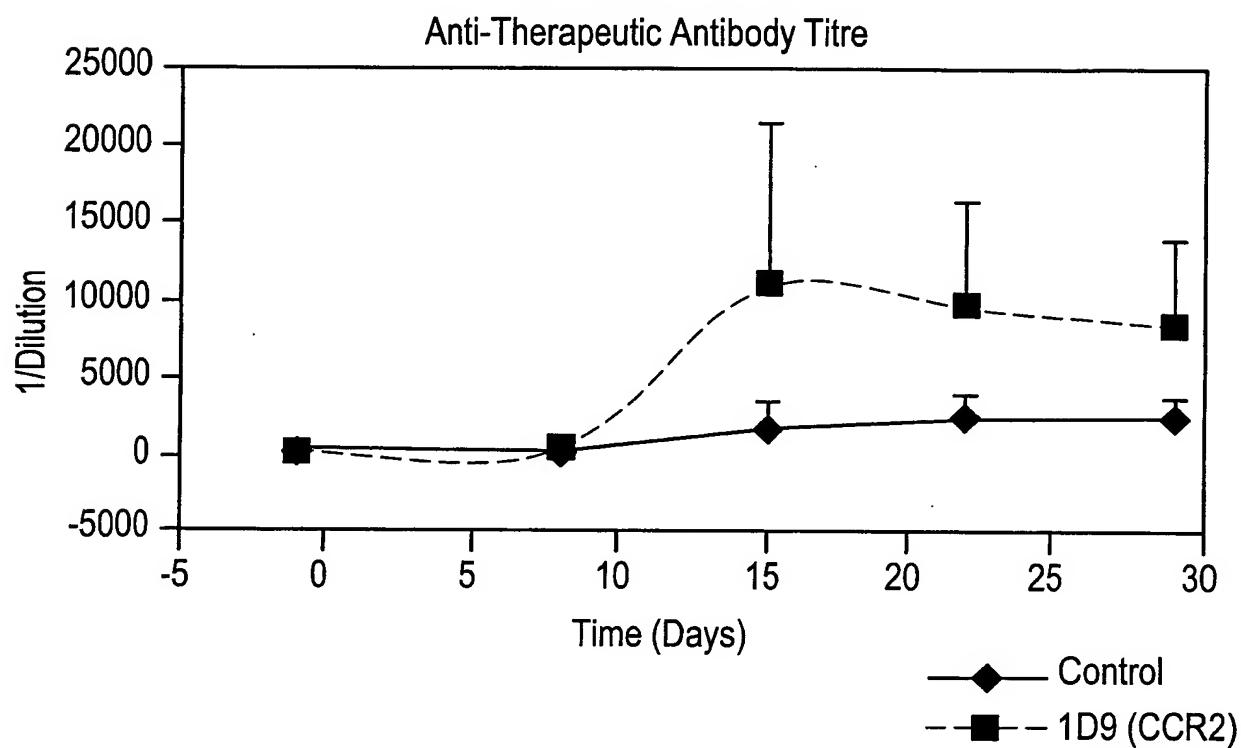
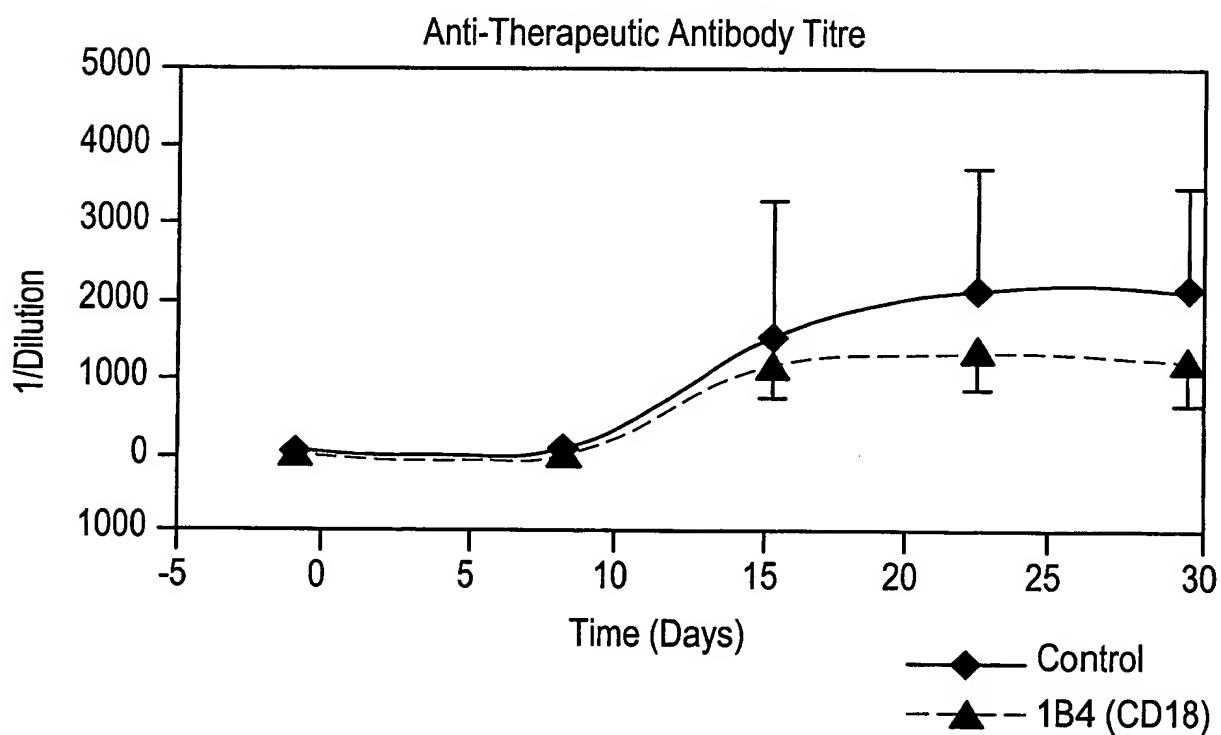


FIG. 4B



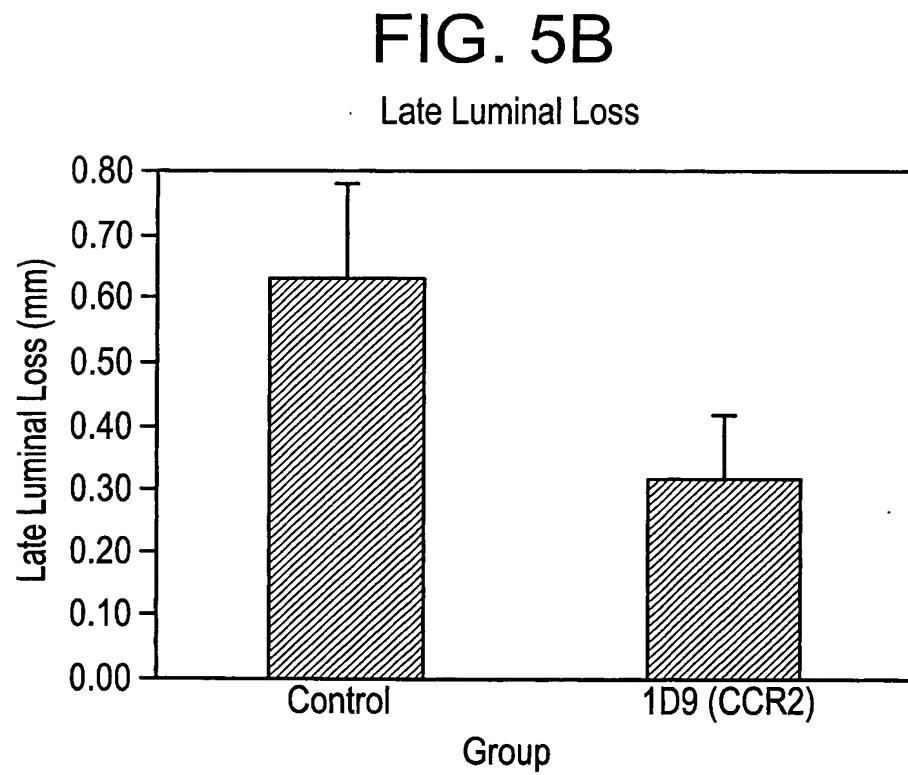
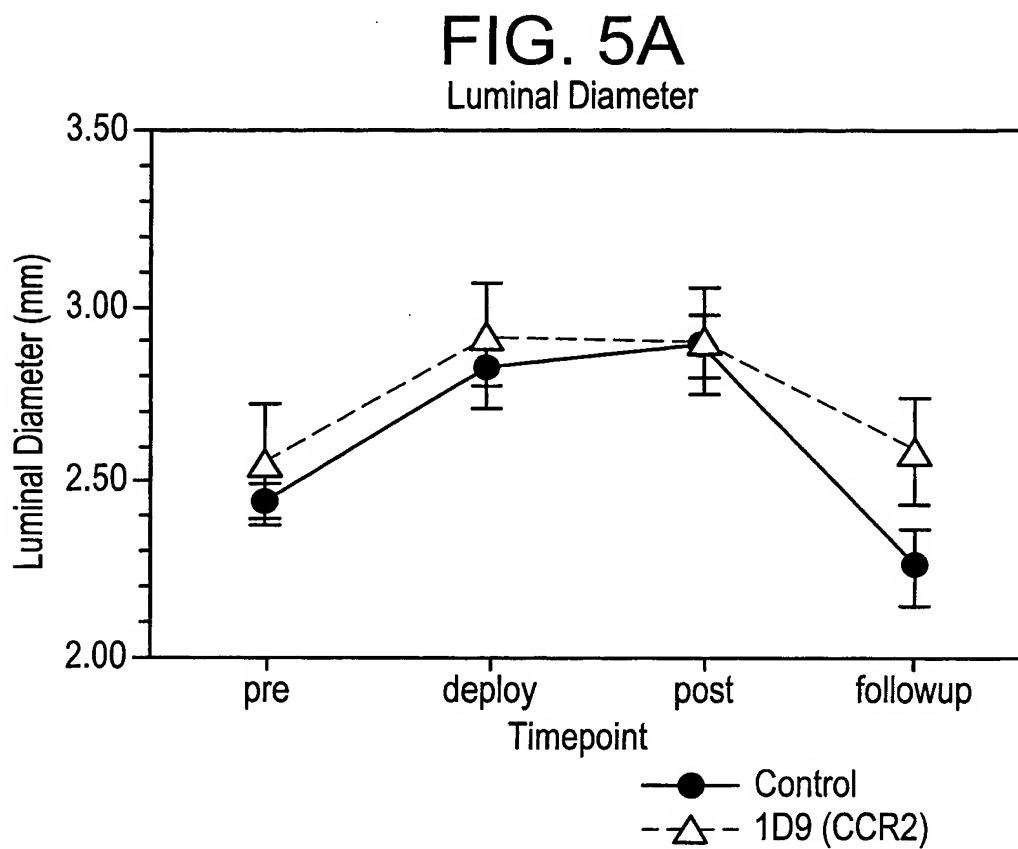


FIG. 5C

Index=LLL/ALG

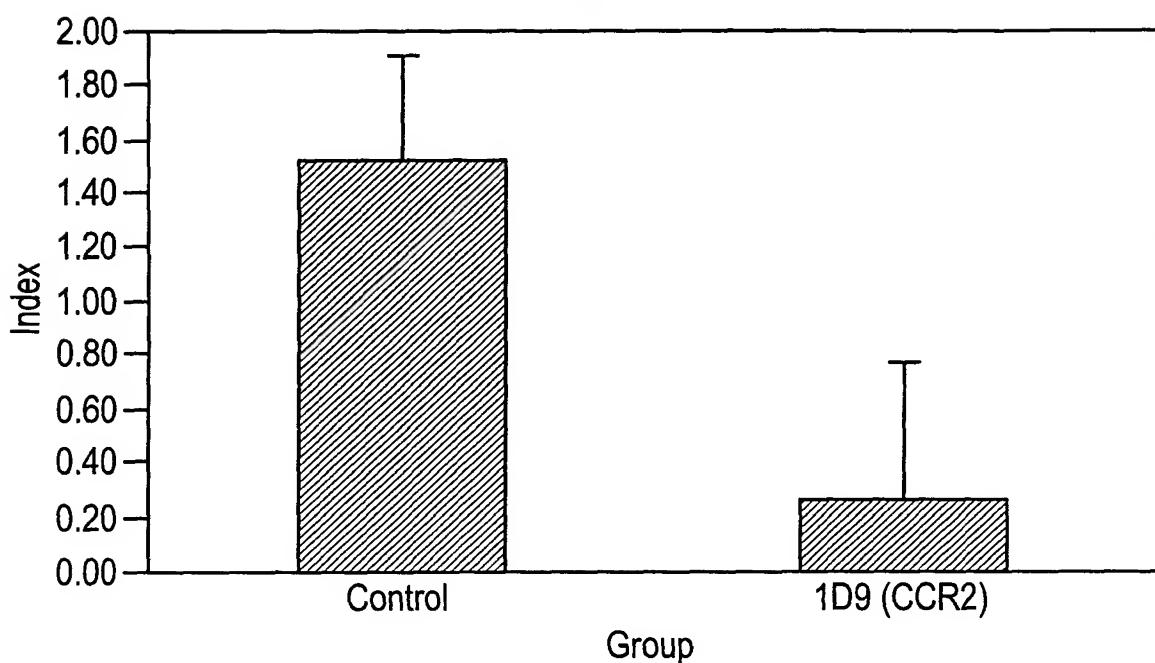


FIG. 5D

Luminal Diameter

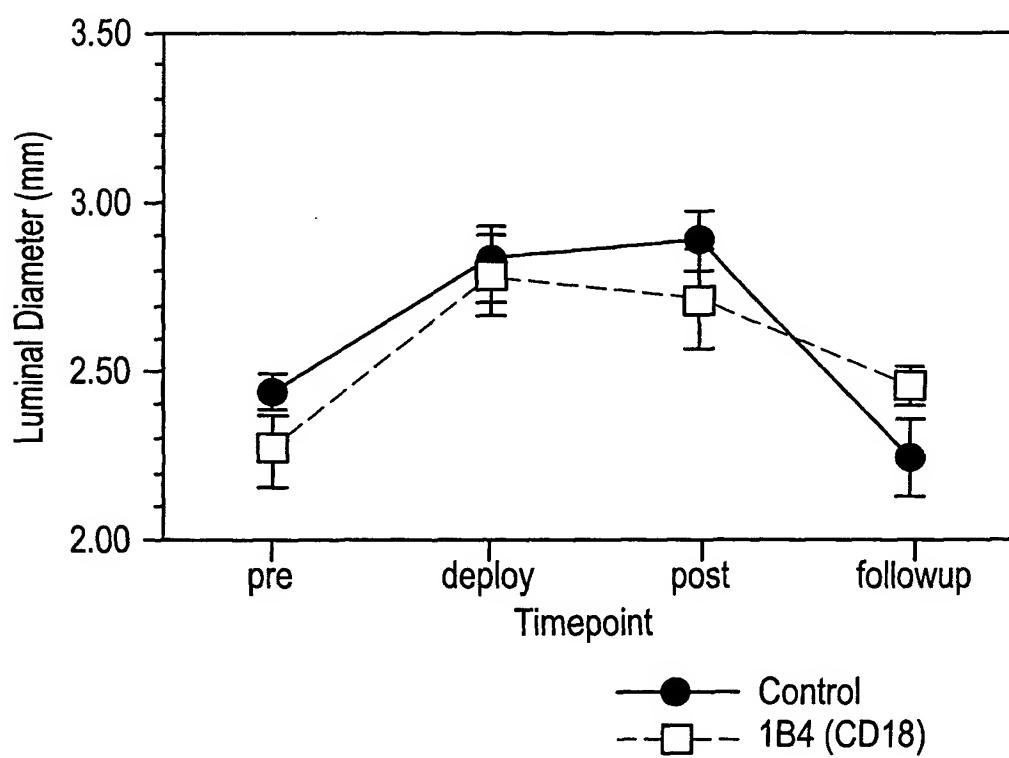


FIG. 5E
Late Luminal Loss

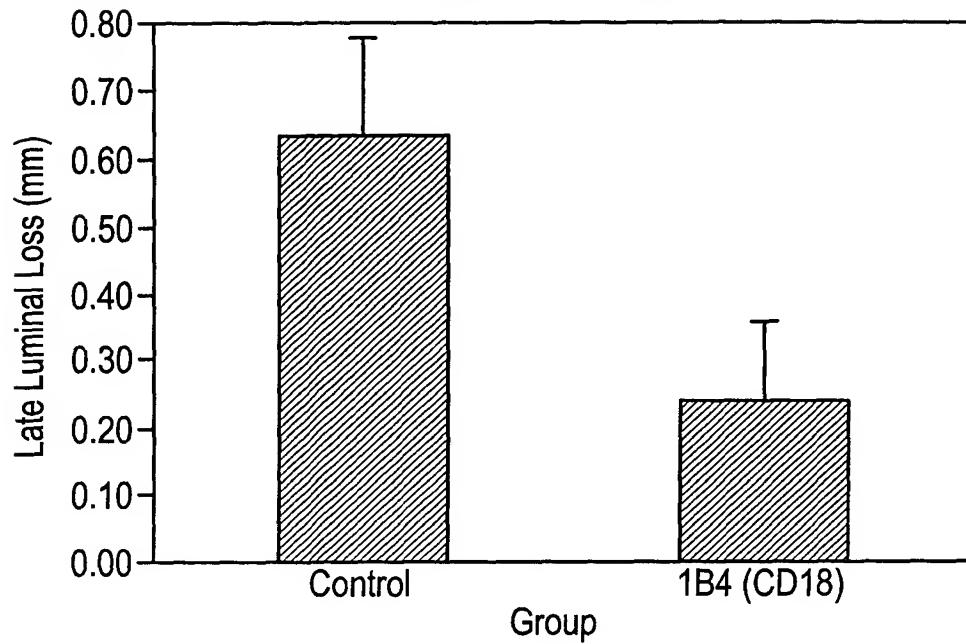


FIG. 5F
Index=LLL/ALG

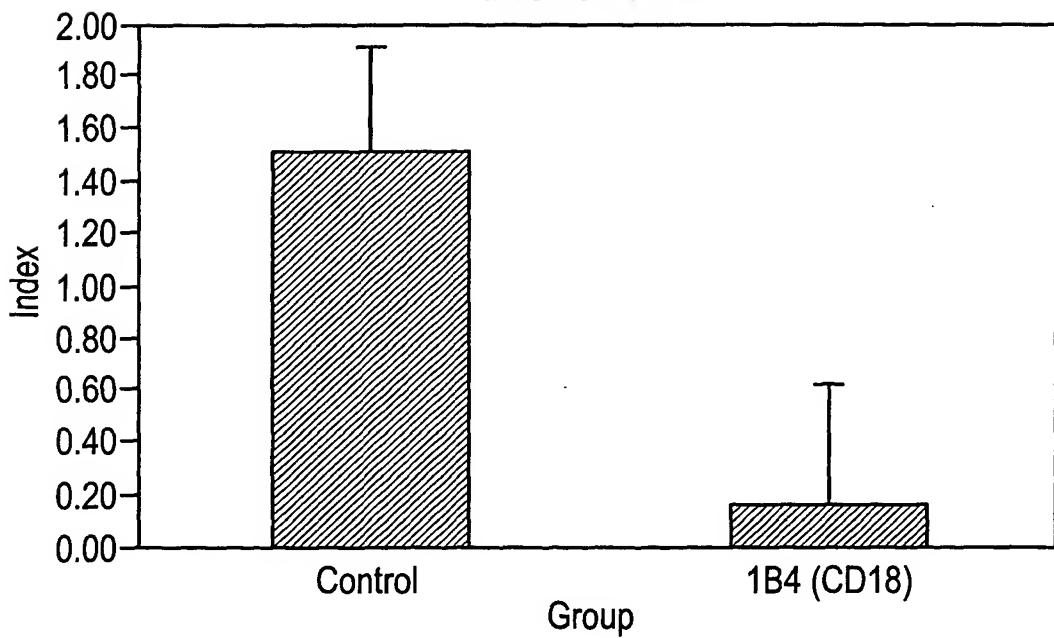


FIG. 6A
Intimal Area (mm²)

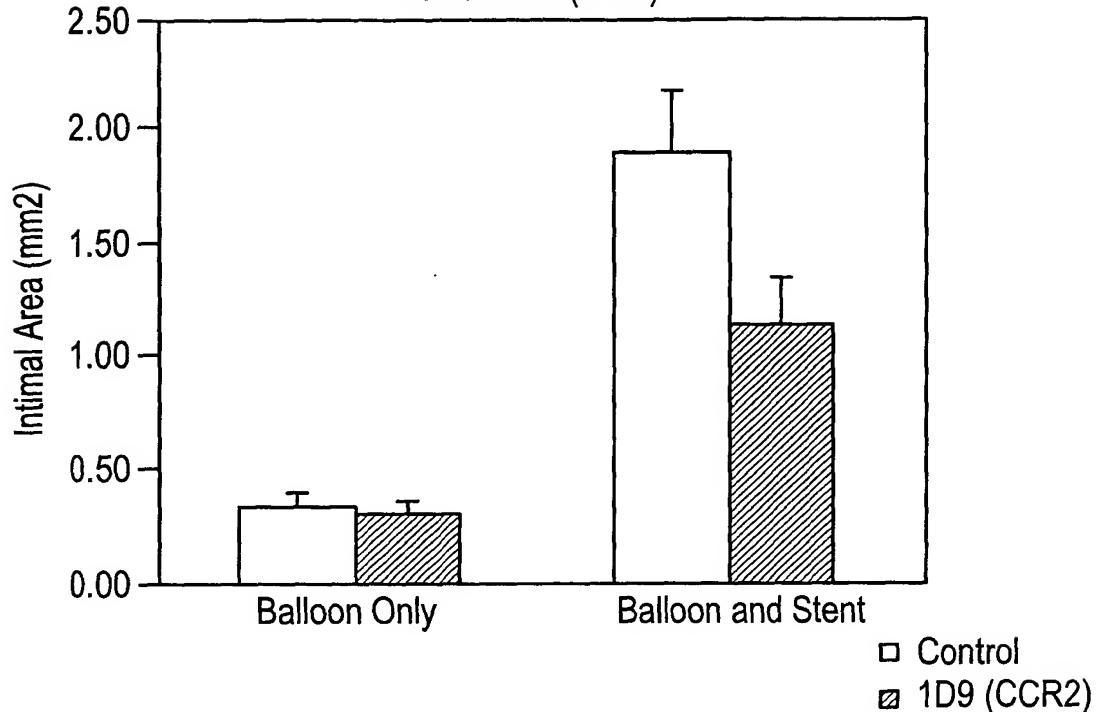
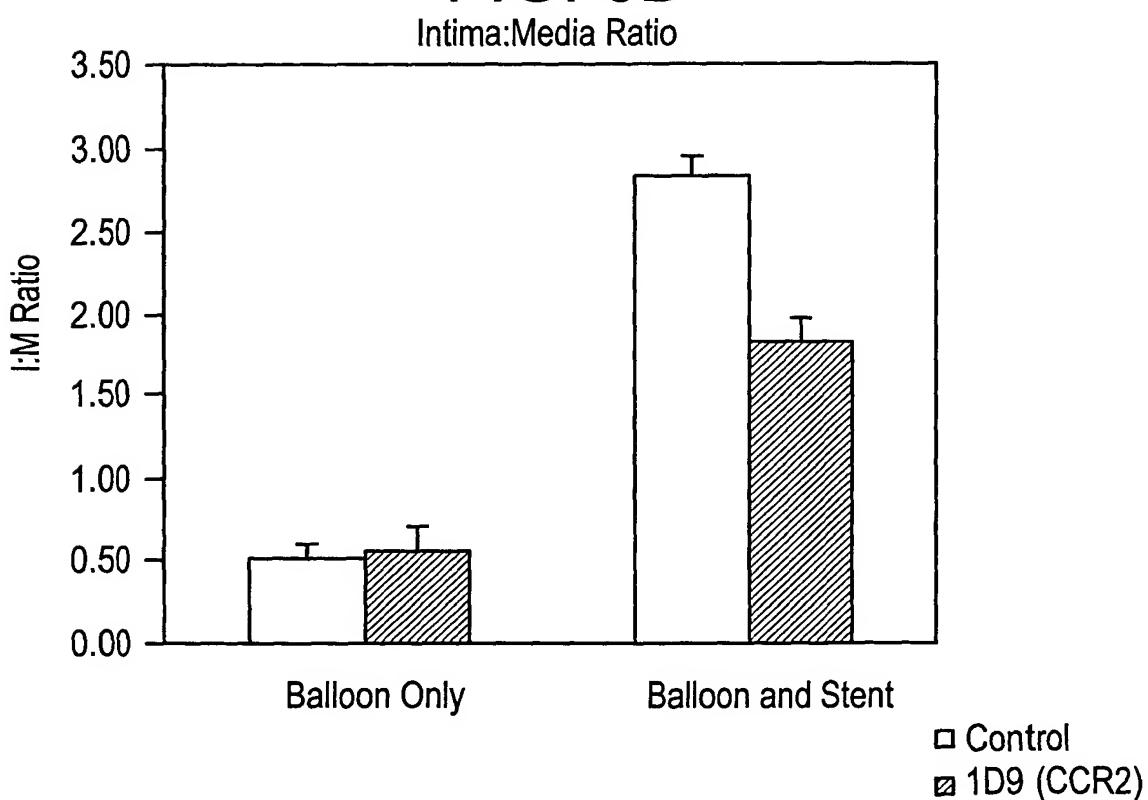
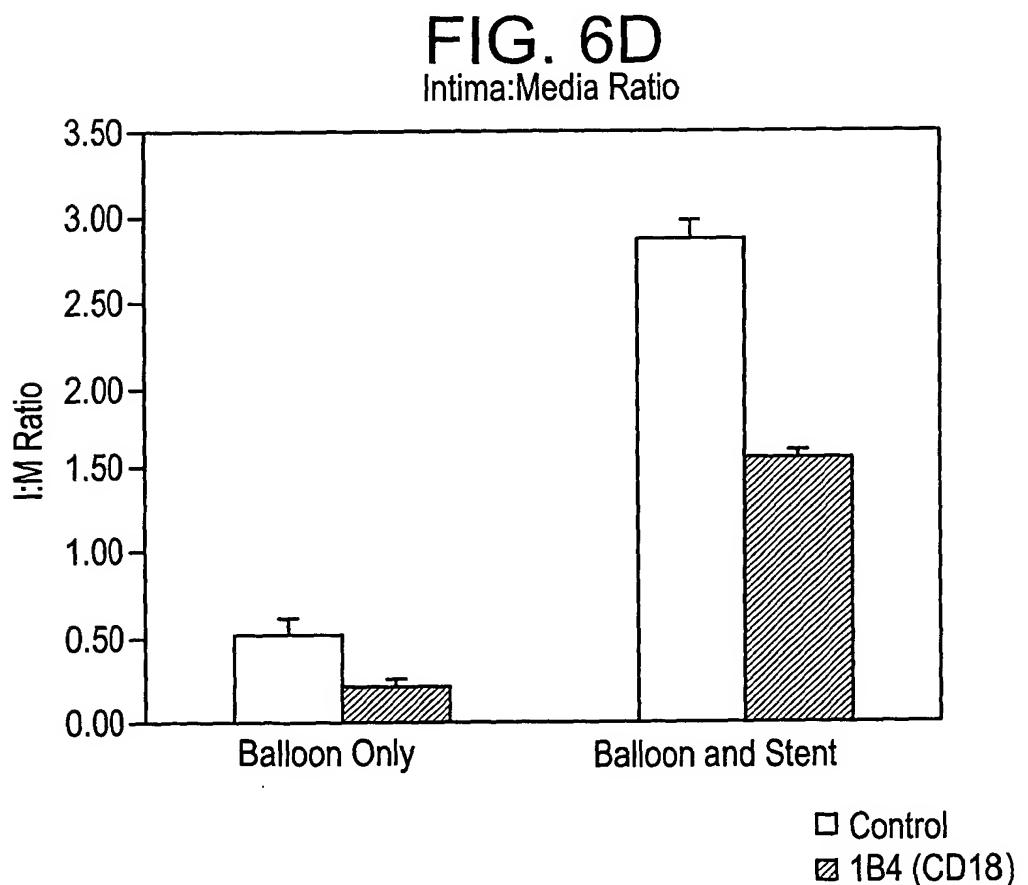
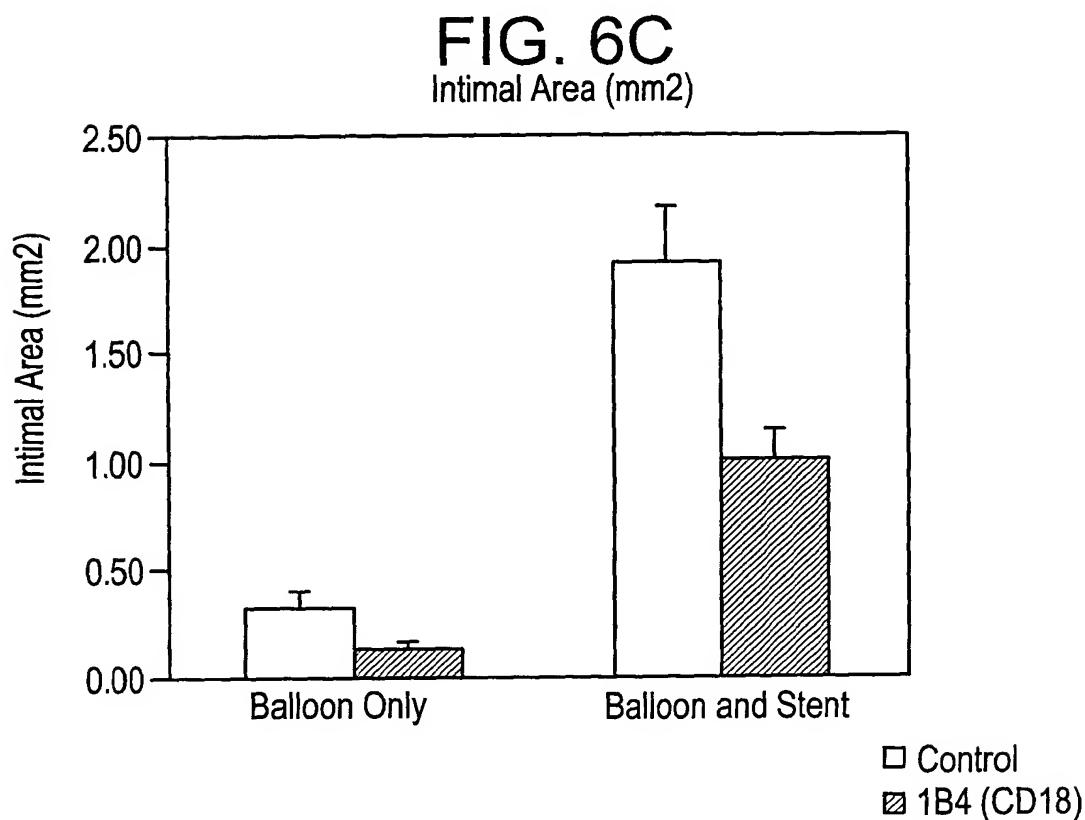


FIG. 6B





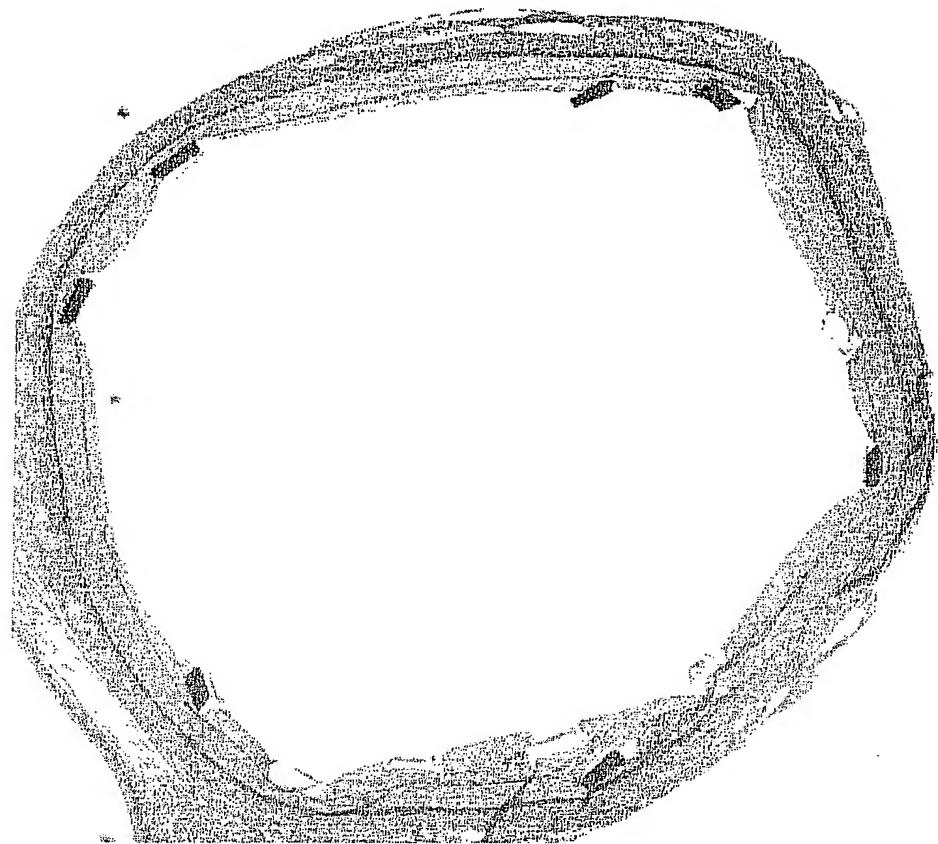


Fig. 7B

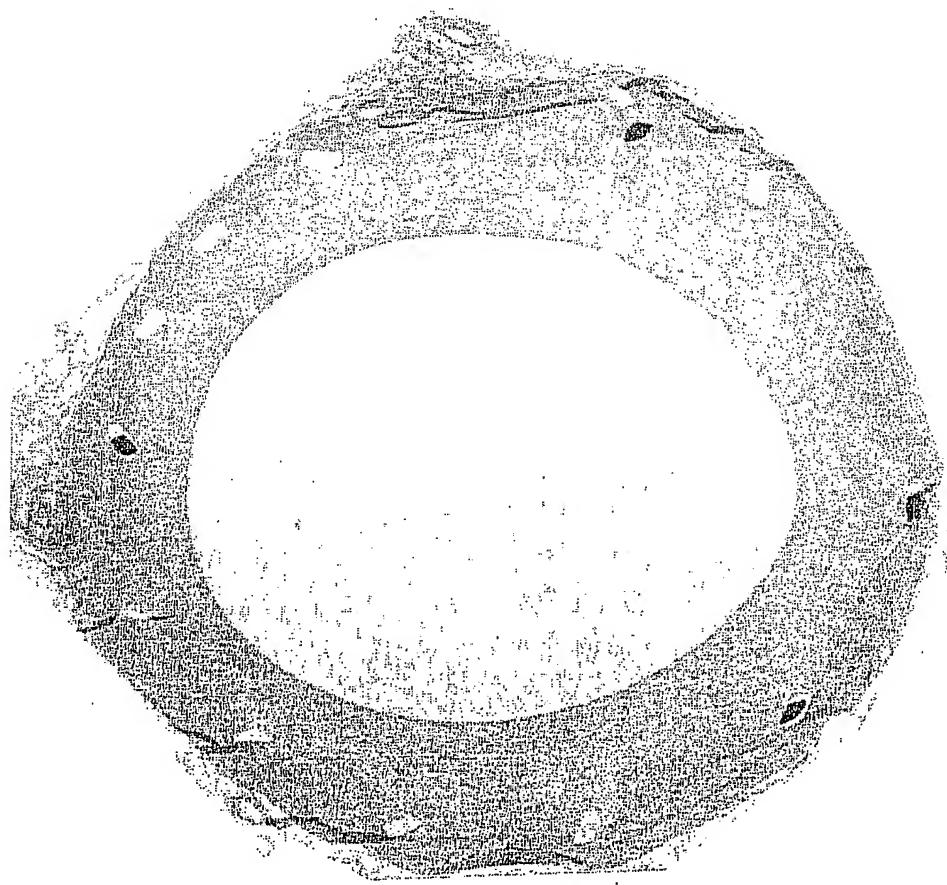


Fig. 7A

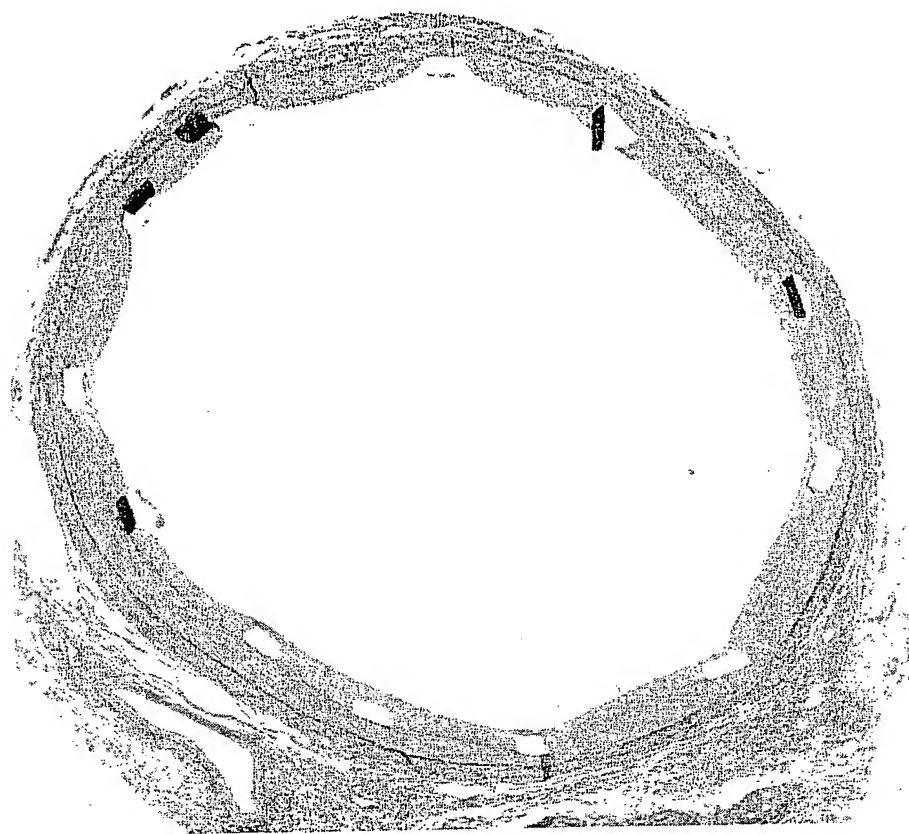


Fig. 8B

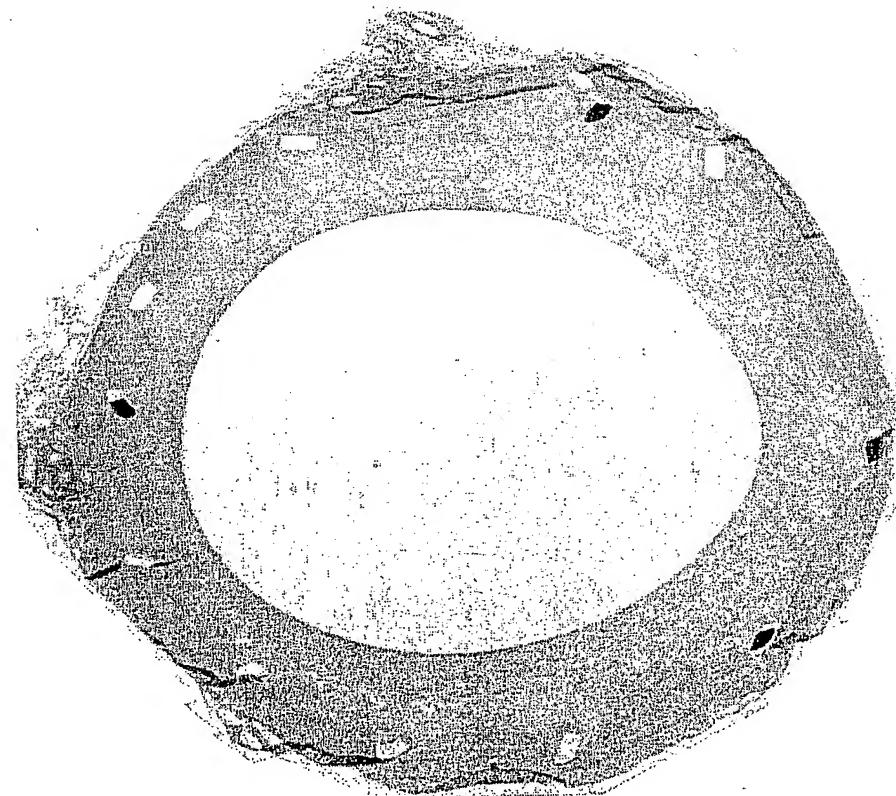


Fig. 8A

Met Arg Val Gln Val Gln Phe Leu Gly Leu Leu Leu Leu Tri Thr Ser

Gly Ala Gln Cys Asp Val Gln Met Thr Gln Ser Pro Ser Tyr Leu Ala

Ala Ser Pro Gly Glu Ser Val Ser Ile Ser Cys Lys Ala Ser Lys Ser

Ile Ser Asn Tyr Leu Ala Tri Tyr Gln Gln Lys Pro Gly Glu Ala Asn

Lys Leu Val Tyr Tyr Gly Ser Thr Leu Arg Ser Gly Ile Pro Ser

Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Arg

Asn Leu Glu Pro Ala Asp Phe Ala Val Tyr Tyr Cys Gln Gln Tyr Tyr

Glu Arg Pro Leu Thr Phe Gly Ser Gly Thr Lys Leu Glu

Fig. 9

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Method of Inhibiting Stenosis and Restenosis
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CDR1	Lys Ala Ser Lys Ser Ile Ser Asn Tyr Leu Ala
CDR2	Tyr Gly Ser Thr Leu Arg Ser
CDR3	Gln Gln Tyr Tyr Glu Arg Pro Leu Thr

Fig. 10

Met Lys Cys Ser Trp Ile Asn Leu Phe Leu Met Ala Leu Ala Ser Gly

Val Tyr Ala Glu Val Gln Leu Gln Ser Gly Pro Glu Leu Arg Arg

Pro Gly Ser Ser Val Lys Leu Ser Cys Lys Thr Ser Gly Tyr Ser Ile

Lys Asp Tyr Leu Leu His Trp Val Lys His Arg Pro Glu Tyr Gly Leu

Glu Trp Ile Gly Trp Ile Asp Pro Glu Asp Gly Glu Thr Lys Tyr Gly

Gln Lys Phe Gln Ser Arg Ala Thr Leu Thr Ala Asp Thr Ser Ser Asn

Thr Ala Tyr Met Gln Leu Ser Ser Leu Thr Ser Asp Asp Thr Ala Thr

Tyr Phe Cys Thr Arg Gly Glu Tyr Arg Tyr Asn Ser Trp Phe Asp Tyr

Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser

Fig. 11

CDR1	Asp Tyr Leu Leu His
CDR2	Trp Ile Asp Pro Glu Asp Gly Glu Thr Lys Tyr Gly Gln Lys Phe Gln Ser
CDR3	Gly Glu Tyr Arg Tyr Asn Ser Trp Phe Asp Tyr

Fig. 12

1 MGWSCHIIFL VATATGVHSQ VQLQESGPGL VRPSQTLSLT CTVSGFTFTD
51 YLLHWVRQPP GRGLEWIGWI DPEDGETKYG QKFQSRVTML VDTSKNQFSL
101 RLSSVTAADT AVYYCARGEY RYNSWFDYWG QGSLVTVSS

Fig. 13

1 MGWSCHILFL VATATGVHSD IOMTOSPSSL SASVGDRVTI TCKASKSISN
51 YLAWYQQKPG KAPKLLIYYG STLRSGVPSR FSGSGSGTDF TFTSSLQPE
101 DIATYYCQQY YERPLTFGQQ TKVEIKR

Fig. 14

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Method of Inhibiting Stenosis and Restenosis
Christopher J. Horvath and Patricia E. Rao

1 DVVMTQTPLT LSVTVGHPAS ISCKSSQSILL DSDGKTFILNW LLQRPQGQSPK
51 RLIYILVSKLD SGVPDRFTGGS GSGTDFTLKI SRVEAEDLGV YYCWQGTHFP
101 YTFGGGTKE IK

Fig. 15

1 EVQLVESGGG LVQPKGSLKL SCAASGFSFN AYAMNWVRQA PGKGLEWVAR
51 IRTKNNNYAT YYADSVKDRY TISRDDSESM LFLQMNNLKT EDTAMYYCVT
101 FYGNGVWGTG TTGTVSS

Fig. 16

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Fig. 17

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4B4'CL	V_h	Chosen human framework acceptor V_h region sequence with mismatches to the 1D9 V_h region highlighted.
1D9RH _A	V_h	CDR grafted 1D9 V_h region, with no back mutations.
1D9RH _B	V_h	CDR grafted 1D9 V_h region, with back mutations at T28S and S30N.
1D9RH _C	V_h	CDR grafted 1D9 V_h region, with back mutations at T28S, S30N, G49A and F67Y.
1D9RH _D	V_h	CDR grafted 1D9 V_h region, with back mutations at T28S, S30N, G49A, F67Y and T93V.

Fig. 18